



## **E. coli infections in poultry - what do we know and where to go.**

**Olsen, Rikke Heidemann; Thøfner, Ida; Ronco, Troels; Pedersen, Karl; Pors, Susanne Elisabeth; Bisgaard, Magne; Christensen, Jens Peter; Kabell, Susanne; Christensen, Henrik**

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# *E. coli* infections in poultry - what do we know and where to go

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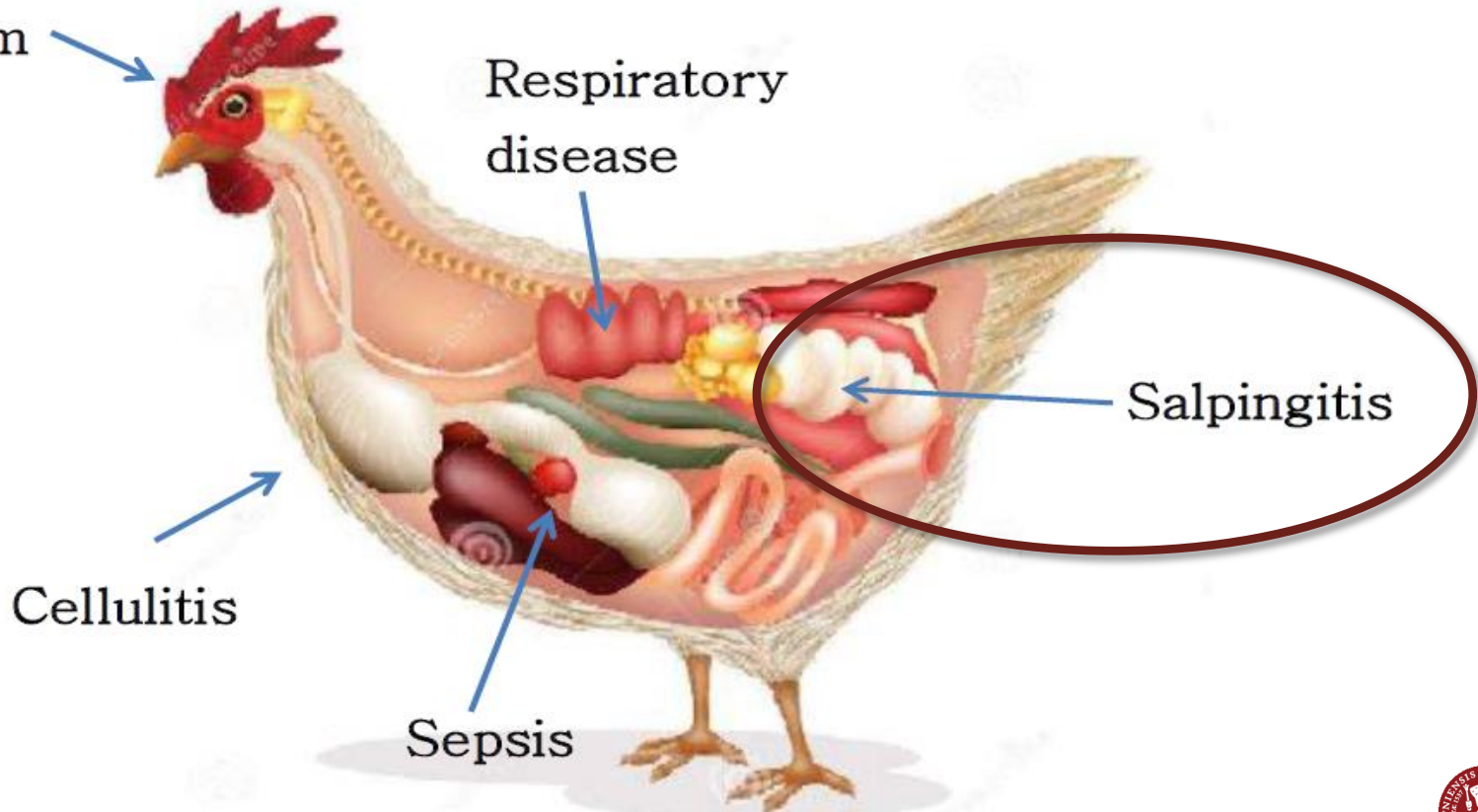
<sup>4</sup> Knowledge Centre for Agriculture, Agro Food Park 15, DK-8200 Aarhus N, Denmark

# Avian Pathogenic *E. coli* (APEC)

Examples of disease manifestations in poultry

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Swollen head  
syndrom



# What characterize an Avian Pathogenic *E. coli* (APEC)?

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- Only focusing *in vitro* on the infections agent (*E. coli*) is not sufficient!
- *In vitro* data must be combined with
  - Flock data (Outbreak/single case?)
  - Host status (Age, vaccination, **viral infection**?)
  - Pathology (Infectious/non infections?)
  - Bacteriology (Other infections, pop. structure)



# Aim

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We aimed to achieve a greater insight into the nature of avian *E. coli* salpingitis combining

- Field outbreak study (bacteriology/pathology)
- Experimental challenge studies
  - Whole genome sequencing analysis of strains of different in vivo pathogenicity



# Materials and methods

## Flock data and bacterial isolation

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- Inclusion criteria:  
Layer flocks with increased mortality due to “colibacillosis”
- Detailed post mortem (pm) examination of submitted layers
- Swabs obtained from liver, spleen and salpinx during pm
- Pure culture *E. coli* isolates further characterized



# Materials and Methods – Field study of outbreak of *E. coli*

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- 8 layers flocks with increased mortality due to *E. coli* (primarily salpingitis)
  - Multilocus sequence typing<sup>1</sup>
  - Phylogenic groups<sup>2</sup>
  - Virulence genes (*astA*, *papC*, *iucD*, *cva/cvi*, *ompT*, *iroN*, F11)<sup>3-5</sup>
- Will not be discussed today due to time

<sup>1</sup> Wirth et al. 2006, <sup>2</sup> Clermont et al. 2000, <sup>3</sup> Ewers et al (2005), <sup>4</sup> Johnson et al. 2006, <sup>5</sup> Johnson et al. 2000



# Results – Overview

| Outbreak number | Sub-mission | Age (wks) | No. of birds | Cause of mortality  | No. of birds demonstrated abundant growth of <i>E. coli</i> in pure culture | Population structure of <i>E. coli</i> |
|-----------------|-------------|-----------|--------------|---|---|--|
| A1              | 1           | 45        | 26           | <i>E. coli</i> (13), cannibalism (13)   | 13 (50%)  | Polyclonal                             |
| B1              | 1           | 41        | 33           | <i>E. coli</i> (23), <i>Streptococcus</i> (8), chronic salpingitis (sterile) accompanied by amyloidosis (1)   | 23 (70%)  | Monoclonal                             |
| C1              | 1           | 78        | 16           | <i>E. coli</i> (11), <i>G. anatis</i> , (2), cannibalism (3)  | 11 (69%)  | Polyclonal                             |
| D1              | 1           | 65        | 56           | <i>E. coli</i> (43), <i>S. aureus</i> (1), cannibalism (11), killed by owner (1)  | 43 (77%)  | Monoclonal                             |
| D2              | 2           | 68        | 48           | <i>E. coli</i> (29), chronic salpingitis accompanied by nephropathy ( <i>E. faecalis</i> ) (1), chronic salpingitis (sterile) accompanied by nephropathy (1), cannibalism (17)                            | 29 (60%)  | Monoclonal                             |
| E1              | 1           | 75        | 42           | <i>E. coli</i> (39), <i>G. anatis</i> , (1), <i>E. faecalis</i> (1), emaciation, dehydration and nephropathy (1)  | 39 (93%)  | Monoclonal                             |
| F1              | 1           | 27        | 12           | <i>E. coli</i> (12)   | 12 (100%)   | Monoclonal                             |
| F2              | 2           | 31        | 16           | <i>E. coli</i> (15), cannibalism (1)  | 15 (94%)  | Monoclonal                             |
| G1              | 1           | 55        | 31           | <i>E. coli</i> (11), cannibalism (6), emaciation, dehydration and nephropathy (1) histomoniosis (1), egg bound (1), amyloidosis(1), liver rupture (1), cellulitis (1), out of lay and killed by owner (2) | 11 (35%)  | Polyclonal                             |
| H1              | 1           | 75        | 43           | <i>E. coli</i> (24), <i>Streptococcus pluranimalium</i> (1), cannibalism (12), uraemia (3), egg bound (3)   | 24 (56%)  | Polyclonal                             |
|                 | 10          |           | 322          |   | 220 (68%)   |  |

Skt. Petersburg, 2017



# Results

## Field study - Multilocus sequence typing

| ST                    | # isolates<br>(n=100) | Prevalence<br>in flocks | Comments  |
|-----------------------|-----------------------|-------------------------|---|
| 23                    | 21                    | 2                       | Mono- and polycl. outbreaks                           |
| 79                    | 4                     | 2                       | Polycl.   |
| 95                    | 33                    | 2                       | Monoclonal  |
| 117                   | 4                     |                         | "Background" monoclonal"                              |
| 140                   | 17                    |                         | Mono- and polycl. outbreaks                           |
| 141                   | 4                     |                         | Polycl.   |
| 357                   | 3                     |                         | Polycl.   |
| 429                   | 4                     |                         | Polycl.   |
| 2490                  | 2                     |                         | Polycl  |
| 67,93,101,155,373,641 | 1/ ST                 | 1-5                     | All except ST101 associated with polyclonal outbreaks |



# Want to know more?



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## **Pathology and Molecular Characterization of *Escherichia Coli* Associated With the Avian Salpingitis-Peritonitis Disease Syndrome**

Author(s): Rikke Heidemann Olsen, Magne Bisgaard, Jens Peter Christensen, Susanne Kabell, and Henrik Christensen

Source: Avian Diseases, 60(1):1-7.

Published By: American Association of Avian Pathologists

DOI: <http://dx.doi.org/10.1637/11237-071715-Reg.1>

URL: <http://www.bioone.org/doi/full/10.1637/11237-071715-Reg.1>

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# Results

## Field study - Multilocus sequence typing

| ST | # isolates<br>(n=100) | Prevalence<br>in flocks | Comments                    |
|----|-----------------------|-------------------------|-----------------------------|
| 23 | 21                    | 2                       | Mono- and polycl. outbreaks |

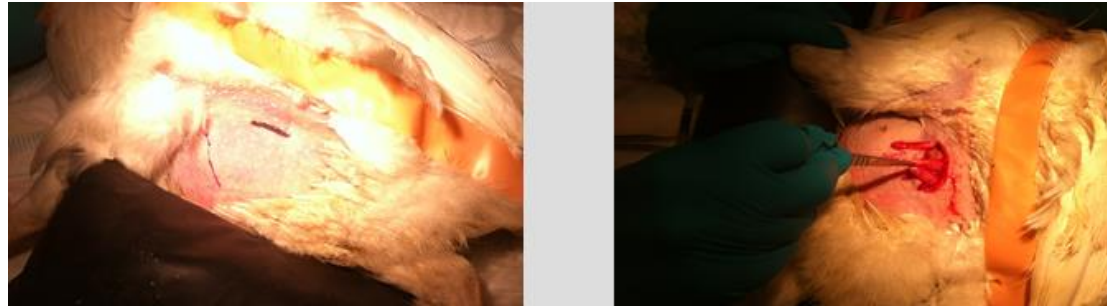
**Model of ascending infection developed;  
Can Koch's postulate be fulfilled?**

|                       |       |     |   |
|-----------------------|-------|-----|---|
|                       |       |     | monoclonal  |
| 140                   | 17    |     | Mono- and polycl. outbreaks                           |
| 141                   | 4     |     | Polycl.   |
| 357                   | 3     |     | Polycl.   |
| 429                   | 4     |     | Polycl.   |
| 2490                  | 2     |     | Polycl  |
| 67,93,101,155,373,641 | 1/ ST | 1-5 | All except ST101 associated with polyclonal outbreaks |



# Materials and Methods – Experimental Challenge Study

- Surgical model<sup>1</sup>
  - 10 Chickens (25 wks)/group
  - Dose 0,1 ml  $6 \times 10^5$  CFU *E. coli*
- Challenge strains



| Strain              | ST  | Phylo-group | Clinical information  |
|---------------------|-----|-------------|---|
| <b>Sept362</b>      | 117 | D           | Isolated from a chicken with septicemia                                       |
| <b>SCI-07</b>       | 369 | B2          | Isolated from a chicken with swollen head syndrom (SHS)                       |
| <b>Chronic_Salp</b> | 429 | B2          | Isolated from a chicken with chronic salpingitis (not outbreak related)       |
| <b>Cp6Salp3</b>     | 95  | B2          | Isolated from a chicken with severe salpingitis and sepsis (outbreak related) |
| <b>D2-2</b>         | 115 | D           | Commensal strain isolated from the cloacae of a healthy chicken               |

- Different infection susceptibility of layers vs. broiler breeders?

<sup>1</sup> Pors et al. 2014.  
Skt. Petersburg, 2017  
Slide 11

## Result and discussion

### Clinical and pathological outcome

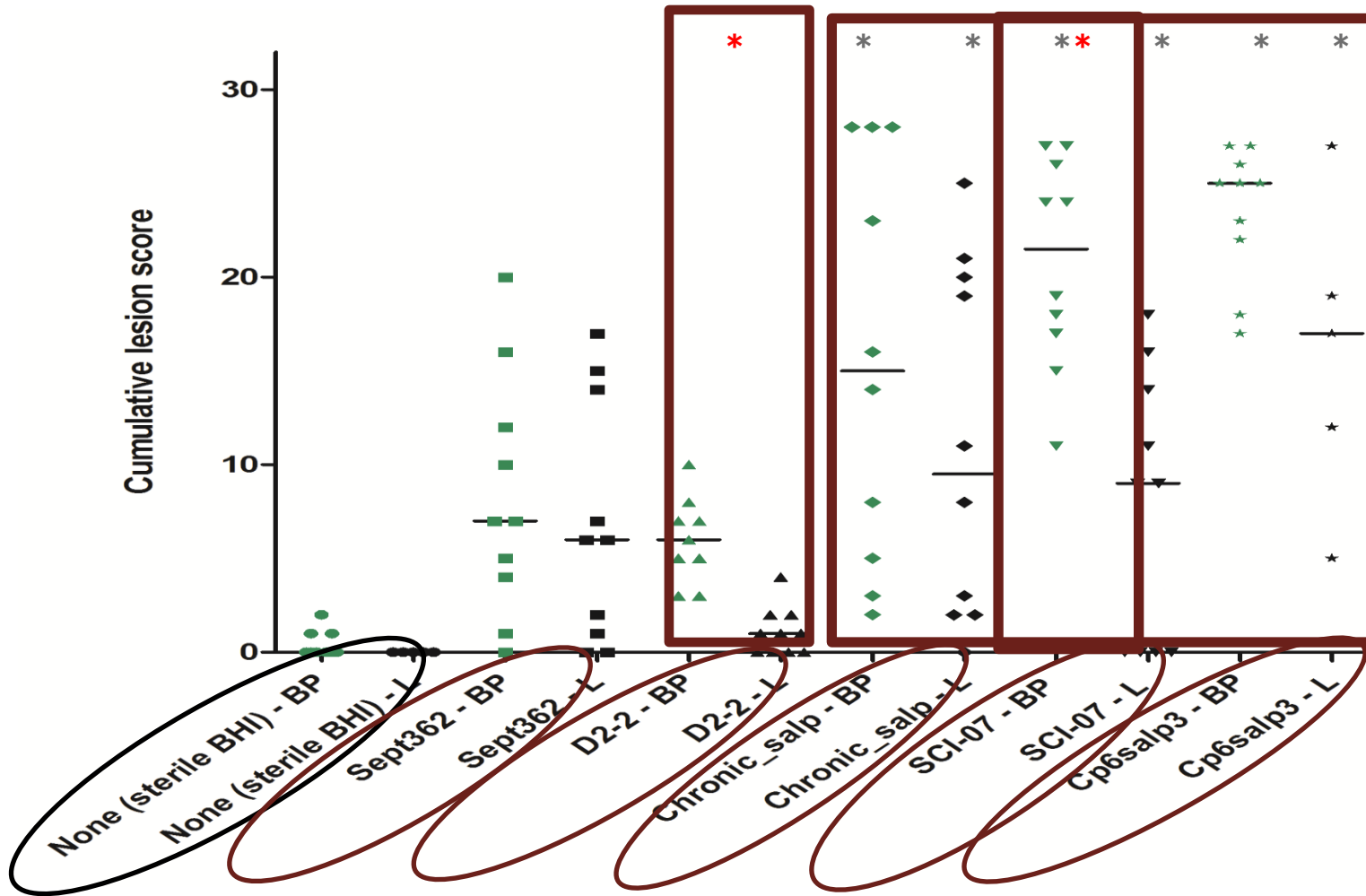
- From limited/none pathology to salpingitis/fibrino-purulent peritonitis/sepsis. Pathology highly similar to field outbreak pathology



# Result and discussion

## Clinical and pathological outcome

\* Sign. Difference between L and BP  
\* Sign. Different from control



## Result and discussion - Genomic comparison of challenge strains

| Characteristics                      | Strains  |  |   |   |  |   |
|--------------------------------------|--|--|---|---|--|---|
|                                      | Cp6Salp3   | Chronic_salp                               | D2-2  | SCI-07  | Sept362  |   |
| Genome size (Mbp)                    | 5,10   | 5,03                                       | 5,68  | 4,97  | 5,28   | NCBI  |
| Sequence type                        | 95   | 429  | 115 <sup>1</sup>                                | 368 <sup>1</sup>  | 117 <sup>1</sup>                                 | <sup>1</sup> MLST 1.7 <sup>12</sup>   |
| Serotype                             | O1:K1:H7   | O83:H4                                     | O2:H9   | H31   | O109:H10   | SeroTypeFinder<br><a href="https://cge.cbs.dtu.dk/services/SeroTypeFinder/">https://cge.cbs.dtu.dk/services/SeroTypeFinder/</a> |
| Plasmids (replicon types)            | IncFII<br>IncFIB   | IncI1, IncFII<br>IncFIB                    | IncI1, <b>IncF</b><br>Col8282, col1156, B/O/K/2 | IncF, IncFII<br>IncFIB, IncFIC,                         | IncI1, IncFII<br>IncFIB, IncFIC, IncFIA          | PlasmidFinder 1.2 <sup>5</sup>  |
| No. predicted pathogen families      | 959*   | 623  | 523   | 477   | 478  | Pathogen finder 1.1 <sup>10</sup>   |
| Virulence, Disease and Defense genes | 109  | 109  | 117   | 124   | 118  | SEED prediction <sup>30</sup>   |
| Iron acquisition and metabolism      | 82   | 53   | 99  | 90  | 80   | SEED prediction <sup>30</sup>   |
| Phage associated genes               | 143*   | 95   | 116   | 79  | 91   | SEED prediction <sup>30</sup>   |
| CRISPR                               | 2  | 1  | 2   | 3   | 2  | CRISPRfinder <sup>12</sup> .  |
| Virulence profile <sup>a</sup>       | Pap operon <sup>b</sup> , vat, upaG, tia, irp2, tsh, iutA, iucA/B/C/D, ehaB, iroN, cva/cvi, maxI, hlyF | irp2, fuyA, tsh, ehaB, cva/cvi, maxI, hlyF | irp2, fuyA, iutA, iucA/B/C/D, iroN, eitB, maxI  | irp2, fuyA, iutA, iucA/B/C/D, iroN, eitB, cva/cvi, maxI | tsh, iutA, iucA/B/C/D, iroN, eitB, cva/cvi, maxI | BRIG <sup>12</sup><br>(Sequences for comparison were obtained at VFDB <sup>12</sup> )   |



# Want to know more?

Veterinary Microbiology 188 (2016) 59–66



Contents lists available at ScienceDirect

Veterinary Microbiology

journal homepage: [www.elsevier.com/locate/vetmic](http://www.elsevier.com/locate/vetmic)



Experimental induced avian *E. coli* salpingitis: Significant impact of strain and host factors on the clinical and pathological outcome



Rikke Heidemann Olsen\*, Ida Cecilie Naundrup Thøfner, Susanne Elisabeth Pors, Teresa Pires dos Santos, Jens Peter Christensen

University of Copenhagen, Department of Veterinary Disease Biology, Faculty of Health and Medical Sciences, Stigboejlen 4, DK-1870 Frederiksberg C, Denmark





# Sum up

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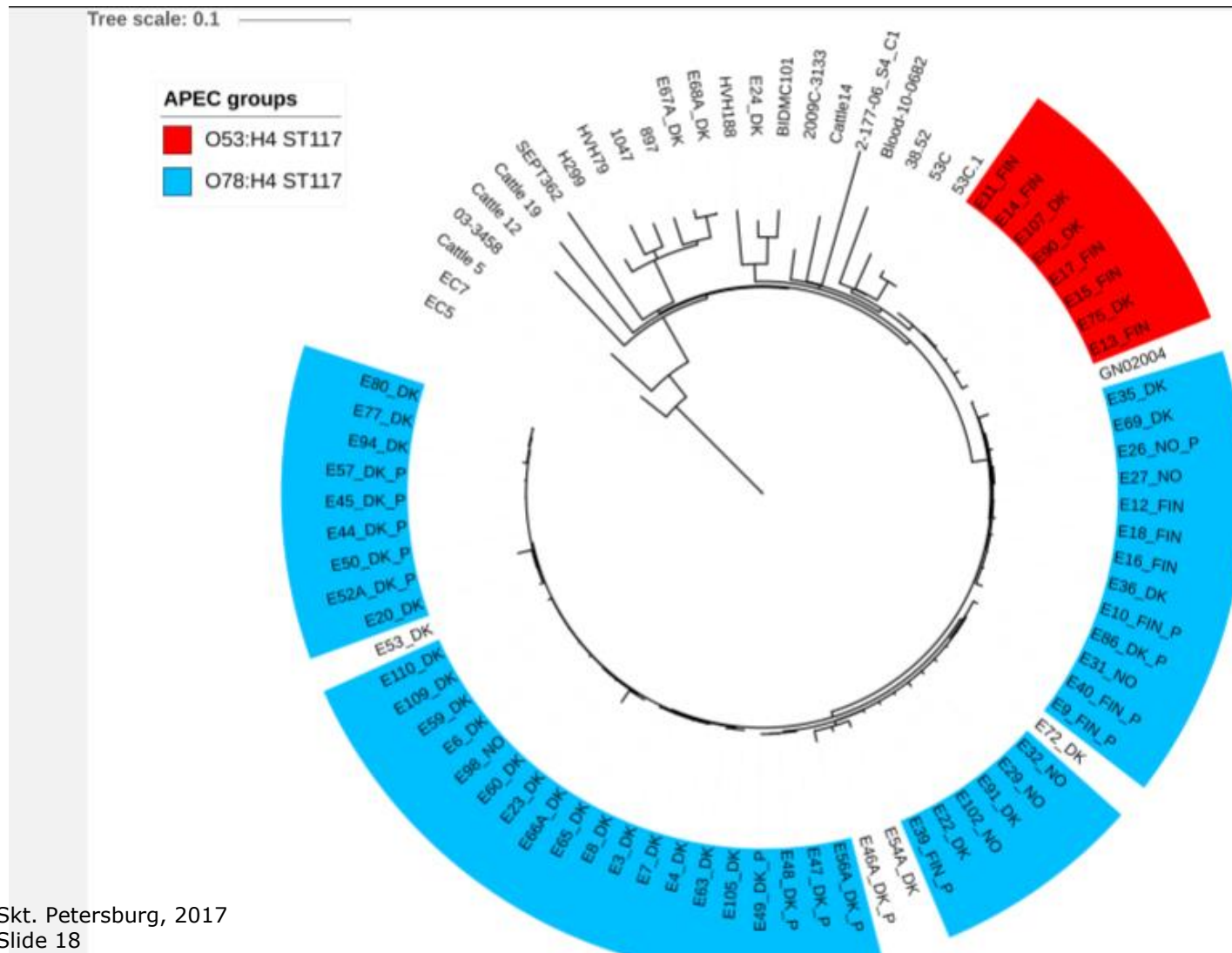
- A broad diversity of *E. coli* strains are capable of causing ascending infection, but is highly depended on **host factors**
- High virulent clones causing **monoclonal outbreaks** in the field
- These high-virulent clones may **represent true APEC**, hence, not being opportunistic
- In general broiler breeders more susceptible than layers



# *E. coli* challenges in the Scandinavian countries 2014-2016



## Whole genome sequencing used to trace the source



# Want to know more?

Ronco et al. *BMC Genomics* (2017) 18:13  
DOI 10.1186/s12864-016-3415-6


BMC Genomics

RESEARCH ARTICLE

Open Access

## Spread of avian pathogenic *Escherichia coli* ST117 O78:H4 in Nordic broiler production



Troels Ronco<sup>1\*</sup> , Marc Stegger<sup>2</sup>, Rikke Heidemann Olsen<sup>3</sup>, Camilla Sekse<sup>4</sup>, Anne Bang Nordstoga<sup>4</sup>, Tarja Pohjanvirta<sup>5</sup>, Berit Lilje<sup>2</sup>, Ulrike Lyhs<sup>1</sup>, Paal Skytt Andersen<sup>2</sup> and Karl Pedersen<sup>1</sup>



# One particular clone of *E. coli* caused the majority of mortality

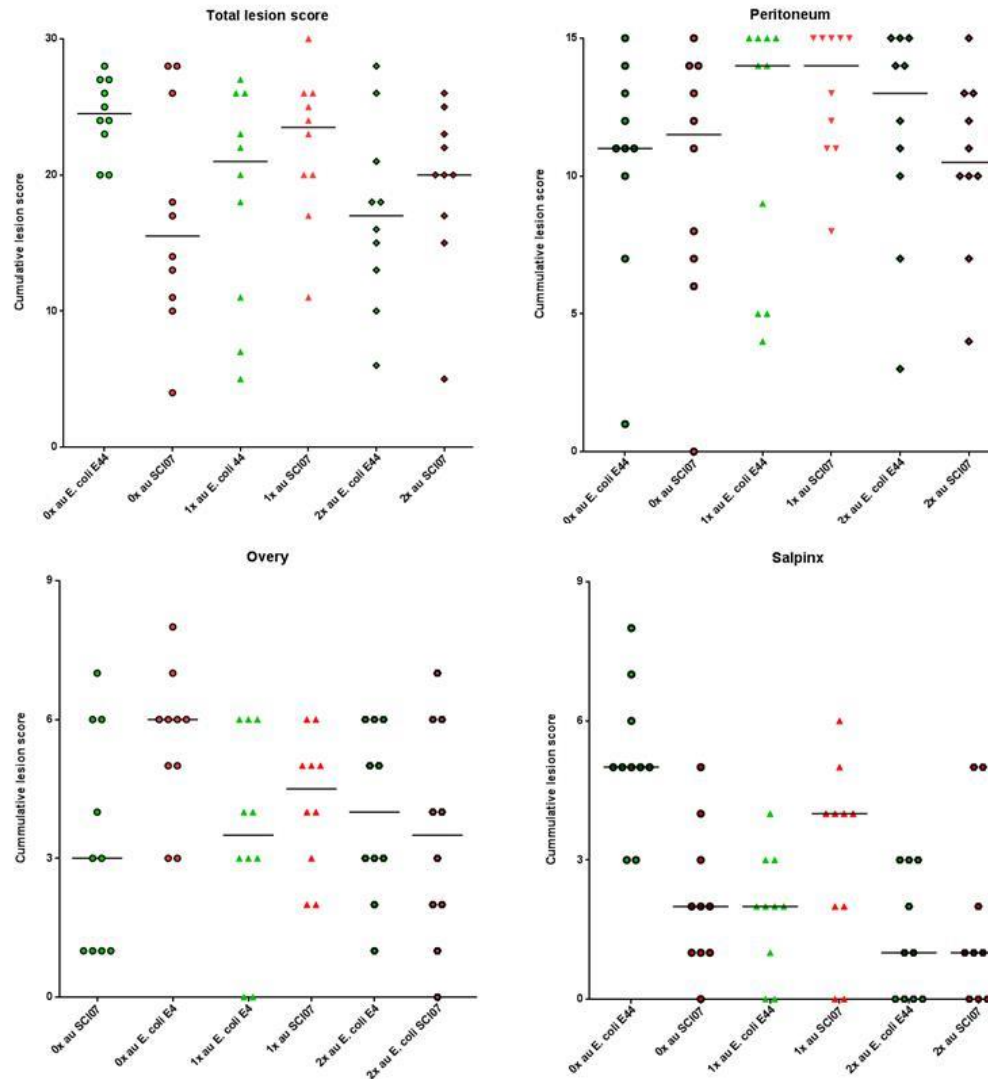
The Danish strategy: Autogenously vaccination  
(+poul vacc)



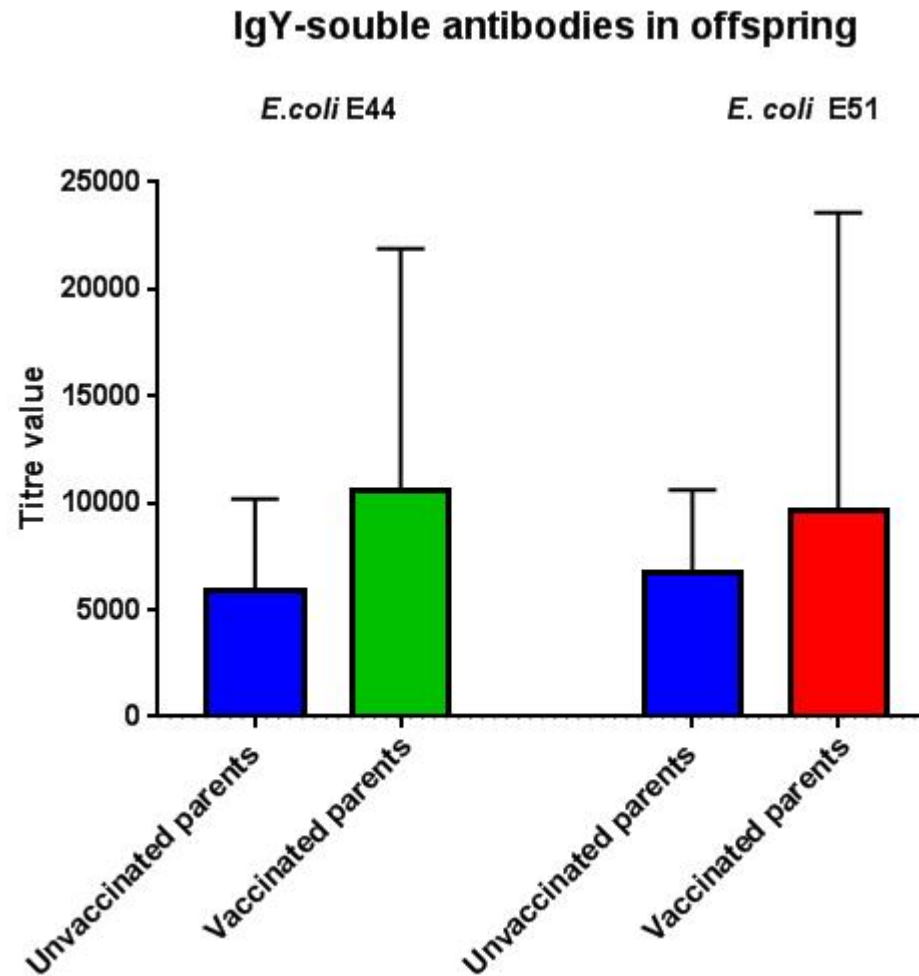
# Experimental set-up to test efficiency of autogenous vaccine

| Group     | E. coli autogenous vaccine (week of age) | Challenge strain <sup>1</sup>             | Average weight at day of challenge | Survived the full experimental period (7 days post challenge) |
|-----------|--|---|------------------------------------|---|
| <b>1A</b> | None                                     | E. coli E44 ( <b>autovaccine strain</b> ) | 3699                               | 0/10 <sup>a</sup>   |
| <b>1B</b> | None                                     | SCI_07                                    | 3669                               | 10/10   |
| <b>2A</b> | 1 time (18)                              | E. coli E44 ( <b>autovaccine strain</b> ) | 3532                               | 8/10  |
| <b>2B</b> | 1 time (18)                              | SCI_07                                    | 3529                               | 10/10   |
| <b>3A</b> | 2 times (14 and 18)                      | E. coli E44 ( <b>autovaccine strain</b> ) | 3563                               | 9/10  |
| <b>3B</b> | 2 times (14 and 18)                      | SCI_07                                    | 3543                               | 10/10   |

# Similar pathology between vaccinated-unvaccinated



# No difference in level of specific antibodies in the offspring





# Want to know more?



Journal

**Avian Pathology** >

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Original Articles

## Evaluation of the efficacy of an autogenous *Escherichia coli* vaccine in broiler breeders

Lili Li, Ida Thøfner, Jens Peter Christensen, Troels Ronco, Karl Pedersen &amp; Rikke H. Olsen

Pages 1-26 | Received 17 Aug 2016, Accepted 25 Nov 2016, Accepted author version posted online: 16 Dec 2016

Download citation

<http://dx.doi.org/10.1080/03079457.2016.1267857>

## Future perspective

- Accept "normal *E. coli* mortality" (Optimize management/biosecurity)
- Focus on the **true APEC**
- The impact of vertical transmission of high-virulent clones of *E. coli* ?
- Target particular clones in a vaccine approach to decrease vertical/horizontal transmission



Thank you for your attention!



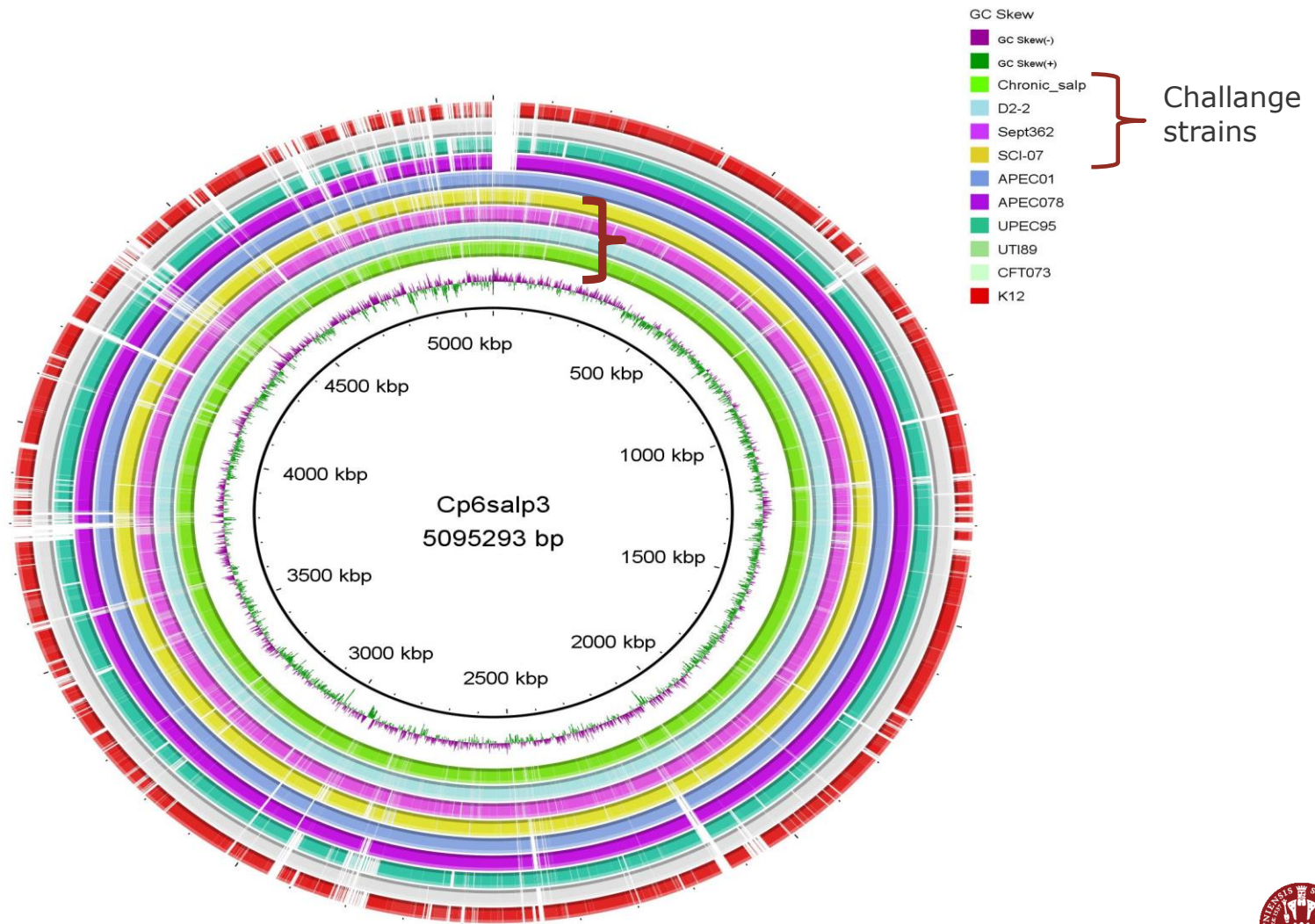
cava@sund.ku.dk



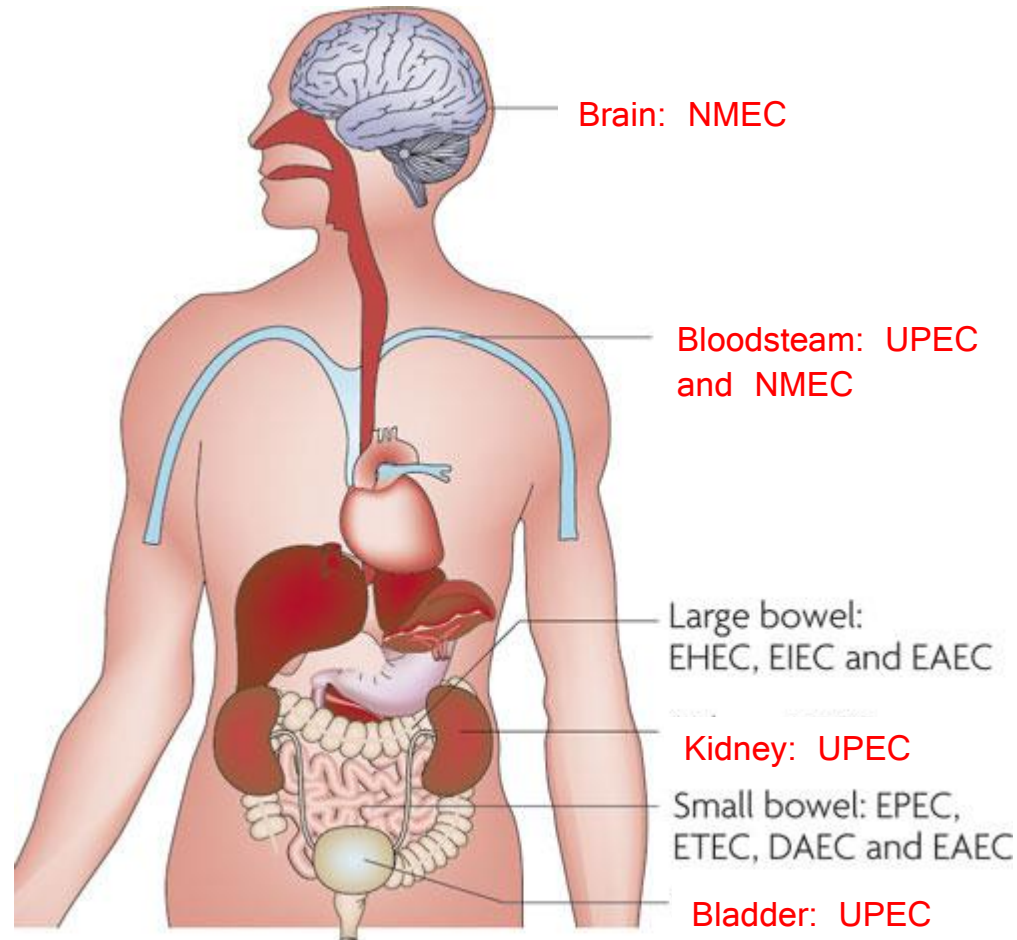
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# Result and discussion

## Genomic comparison of challenge strains



# Pathotypes of human pathogenic *E.coli*



Croxen and Finlay (2010)

Nature Reviews | Microbiology

# Definition of an *E. coli* pathotype (In human medicine)

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“A defined pathological lesion caused by a population of *E. coli* having characteristic mechanisms to attach to and exploit the host's cells of a particular niche”



